

عنوان مقاله:

Genotype-environment interaction and stability analysis for seed yield of Fenugreek (*Trigonella foenum-graecum* L.) genotypes

محل انتشار:

ماهنامه پیشرفت های کشاورزی، دوره 9، شماره 7 (سال: 1399)

تعداد صفحات اصل مقاله: 6

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خلاصه مقاله:

Yield data of 12 Fenugreek (*Trigonella foenum-graecum* L.) genotypes tested across 9 rain-fed environments during the 2015-2017 growing season using RCBD in 3 replications were analyzed using the AMMI model. The AMMI analysis tested in nine environments (years) were showed that the yield was significantly affected ($P < 0.001$) by environment main effects. However, it was not significantly affected by genotypes main effect as well as GxE interaction. The model revealed that differences between the environments accounted for about 91.28% of the treatment sum of squares. The genotypes and the GxE interaction also accounted significantly for 0.92% and 7.79% respectively of the treatment SS. The first principal component axis (PCA 1) of the interaction captured 48.61% of the interaction sum of squares. Similarly, the second principal component axis (PCA2) explained a further 24% of the GEI sum of squares. The mean squares for the PCA 1 and PCA 2 were significant at $P = 0.01$ and cumulatively contributed to 72.61% of the GxE interaction SS, leaving 27.44% of the variation in the GxE interaction in the residual. The AMMI and AMMI stability value (ASV) identified G7 and G5 as the stable and high yielding genotypes. Yield data of 12 Fenugreek (*Trigonella foenum-graecum* L.) genotypes tested across 9 rain-fed environments during the 2015-2017 growing season using RCBD in 3 replications were analyzed using the AMMI model. The AMMI analysis tested in nine environments (years) were showed that the yield was significantly affected ($P < 0.001$) by environment main effects. However, it was not significantly affected by genotypes main effect as well as GxE interaction. The model revealed that differences between the environments accounted for about 91.28% of the treatment sum of squares. The genotypes and the GxE interaction also accounted significantly for 0.92% and 7.79% respectively of the treatment SS. The first principal component axis (PCA 1) of the interaction captured 48.61% of the interaction sum of squares. Similarly, the second principal component axis (PCA2) explained a further 24% of the GEI sum of squares. The mean squares for the PCA 1 and PCA 2 were significant at $P = 0.01$ and cumulatively contributed to 72.61% of the GxE interaction SS, leaving 27.44% of the variation in the GxE interaction in the residual. The AMMI and AMMI stability value (ASV) identified G7 and G5 as the stable and high yielding genotypes.

کلمات کلیدی:

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